

AGCAGAGAGCCTGGTGGGCATGGACATCTTTATCCACATACCTTAGTGTGAC  
 CACGCCGACAGAAACTACTAAGGCCATCTCAGGGGTGCCTGTGCCAGGAGA  
 GGGGGGCGGTGTCCCCGGGCCGAGAGCCATGCCTTTCGGCCTGAAGCTCCG  
 CAGGACTCGGCGCTACAACGTCCTGAGCAAGAACTGCTTTGTTGCCCGGATC  
 CGCCTGCTGGACAGCAATGTCATCGAGTGCACGCTGTCGGTGGAAGCACGG  
 GGCAAGAGTGCCTGGAGGCCGTGGCCCAGAGGCTGGAGCTGAGGGAGACGC  
 ACTACTTCGGCCTTTGGTTTCTCAGCAAGAGCCAGCAGGCGAGATGGGTAGA  
 GCTGGAGAAGCCACTGAAGAAACATCTGGACAAGTTTGCTAACGAGCCTCTG  
 CTTTCTTCGGAGTCATGTTCTATGTGCCAAATGTGTACGGCTTCAGCAGGA  
 GGCCACAAGATATCAGTATTACCTGCAAGTCAAAAAGACGTGCTTGAAGGA  
 CGGTTGCGGTGCTCCCTGGAACAAGTGATCCGGCTGGCTGGCTTAGCTGTGC  
 AAGCTGACTTCGGAGATTATAACCAGTTTGATTCCCAAGAGTTCCTCCGAGA  
 GTATGTGCTCTTTCCTATGGATTTGGCCATGGAGGAGGCGGCTCTGGAGGAG  
 CTAACCCAGAAGGTGGCCCAGGAACACAAAGCTCATAGCGGGATCCTGCCG  
 GCTGAAGCTGAACTGATGTACATCAACGAGGTAGAGCGTTTGATGGATTTG  
 GACAGGAGATCTTCCCCGTGAAGGACAGTCATGGCAACAGCGTGCACCTCGG  
 CATCTTCTTCATGGGGATTTTTGTGAGGAACAGGGTCGGGAGACAGGCAGTG  
 ATATACAGGTGGAATGACATTGGGAGTGTTACTCACAGCAAAGCAGCCATCC  
 TGTTGGAGCTGATTGACAAGGAGGAGACCGCGCTCTTCCATACAGATGATAT  
 TGAAAATGCCAAGTACATTTCTCGGTTGTTTACCACTCGGCACAAATTTTACA  
 AACAGAACAAGATCTGCACTGAACAGTCAAATTCTCCACCCCCAATCAGACG  
 CCAGCCCACCTGGAGCCGGTCTCACTGCCAAGGCAGCAGCCGTATATCTTG  
 CCTCCCATGCATGTCCAGTGCAGTGAGCACTACTCGGAGACCCATACTTCCCA  
 AGACAGCATTTTCCCCGGGAACGAAGAAGCCTTGTACTGCCGTTCTCACAAC  
 AGCCTGGACCTTAATTACTTGAACGGCACCGTCAACCAATGGCAGCGTGTGCA  
 GCGTTCACAGCGTCAACTCCCTCAGCTGCTCCAGAGCTTCATTCAGGCGTCT  
 CCAGTGTCTTCCAACCTTAGCATCCCTGGGAGTGACATCATGAGGGCCGATT  
 ACATCCCCAGCCACCGCCACAGCACCATCATCGTGCCGTCTTACAGGCCGAC  
 CCCAGATTACGAGACGGTCATGAGGCAGATGAAGAGGGGTCTGATGCACGC  
 AGACAGCCAGAGCCGGTCTCTGCGTAACCTCAATATCATCAACACCCATGCC  
 TATAACCAGCCCGAGGAACTGGTGTACAGCCAGCCGGAGATGCGGGAGAGG  
 CATCCCTACACGGTCCCCTATGCACACCAGGGGTGCTACGGTCACAACTTG  
 TAAGTCCGTCTGACCAGATGAACCCCCAAAATTGTGCGATGCCTATCAAGCC  
 AGGGGCCAGTTCCATCTCTCACACAGTGAGCACTCCAGAACTAGCCAACATG  
 CAGCTCCAAGGAGCACAACACTATAGCACAGCCACATGCTCAAGAACTATC  
 TATTCAGGCCGCCACCCCCCTTACCCTCGGCCCCGTCCTGCCACCAGCACCCCA  
 GACCTCGCCAGCCACCGCCACAAGTACGTACGCGGCAGCAGCCCTGATCTGG  
 TAACTCGGAAGGTGCAGCTCTCCGTAAAGACCTTCCAGGAGGACAGCTCACC  
 TGTGGTCCATCAGTCTCTGCAGGAGGTGAGCGAACCCTCACAGCCACCAAG  
 CACCATGGCGGGCGGGTGGCACGGTGAATAAACGCCACAGCCTGGAGGTG

FIGURE 1A

ATGAACAGCATGGTGAGAGGCATGGAGGCCATGACACTGAAGTCACTCAATA  
TCCCCATGGCTCGCCGCAACACCCTTCGGGAGCAGGGCCCTTCCGAGGAGAC  
GGGCGGCCACGAAGTGCACGGTCTCCCCCAGTATCACCACAAGAAGACATTC  
TCGGATGCCACCATGCTGATCCACAGCAGTGAGAGCGAGGAAGAGGAGGAG  
ACCCTGGAGGCTGCACCTCAGGTTCTGTGCTTCGAGAGAAAGTAGAATACA  
GTGCCCAGCTGCAGGCTGCCCTGGCCCGCATCCCCAACAGGCCCCCACCTGA  
GTACCCAGGGCCAAGAAAAAGTGTGAGTAATGGGGCACTGAGACAGGACCA  
GGGAACCCCTCTTCCTGCCATGGCCAGGTGCAGGGTGCTGAGACACGGACCA  
TCCAAGGCCCTCAGTGTCTCCCGGGCAGAGCAGCTGGCTGTCAACGGTGCCT  
CTCTGGGTCCCTCCATCTCTGAGCCTGACCTAACCAGCGTGAAGGAGCGGGT  
CAAGAAAGAGCCTGTGAAGGAAAGGCCGGTGTGAGAGATGTTCTCCCTGGAG  
GACAGCATTATAGAGAGAGAGATGATGATCAGGAATCTAGAGAAGCAGAAG  
ATGACGGGCCCCGCAGGCACAGAAGAGACCGCTGATGTTGGCAGCGCTGAAT  
GGGCTCTCGGTGGCCCGAGTGTGCGGGGCGGGAAGATGGTCGCCATGATGCCA  
CCCGAGTCCCCATAGACGAGAGGCTCAGAGCCCTGAAGAAGAAGCTGGAAG  
ATGGAATGGTGTTCACAGAATATGAGCAGATTCCAAACAAAAAGGCCAACG  
GCGTCTTCAGCACCGCCACTCTGCCTGAGAACGCCGAGCGCAGCCGGATCCG  
AGAAGTTGTCCCATATGAGGAGAATCGAGTGGAGCTCATCCCGACCAAAGAA  
AACAACACAGGCTATATCAACGCCTCCACATCAAGGTGGTGGTCGGCGGAT  
CAGAATGGCACTACATCGCCACCCAGGGGCCCTTGCCACATACGTGCCATGA  
CTTCTGGCAGATGGTGTGGGAGCAGGGGGTGAATGTGATCGCCATGGTCACT  
GCAGAGGAGGAGGGTGGACGGACCAAAAGCCATCGATACTGGCCCAAAGT  
GGGTCCAAGCATAGTTCTGCCACCTACGGCAAGTTCAAGGTCAACACAAAGT  
TCCGGACAGATTCTGGTTGCTATGCAACGACGGGCCTAAAGGTGAAGCACCT  
GCTGTCCGGGCAGGAGAGGACCGTGTGGCACTTGCAGTACACGGAAGTGGCCC  
CACCACGGCTGTCCAGAAGACGTCCAAGGATTTTTGTCTACTTGGAGGAAA  
TCCAGTCAGTCCGACGCCACACCAACAGCGTGCTGGAAGGCATCAGGACCAG  
GCACCCCCCATCGTGGTTCACTGCAGCGCGGGTGTGGGAAGGACTGGTGTG  
GTTATCCTCTCTGAGCTCATGATCTACTGCCTGGAACACAACGAAAAGGTGG  
AGGTGCCCACGATGCTGCGATTCTCAGGGAGCAGAGGATGTTTCATGATCCA  
GACCATTCGCGAGTACAAGTTCGTCTACCAAGTCCTCGTCCAGTTCTGCAGA  
ATTCCAGGCTCATTTGATCTCCTCCGGGATGCAGCTTCTGGAGGAGGGACGC  
AGCTCTGTCCTGCAGGGGGCGGCCACTTCGACAACATCTGCCTCCCCCAGCC  
AGAGGTGGATGGCTGGCAGCAGGCAGAAGCCAGAGTTACTCACAAACATCA  
TGTATTATTTTATATAAGATAATTTATTTTTTCCCTCTTTGGAATAAGTTCTG  
TGAGTTATTATATAATGCTTCCCCCCCATACACACACAATAATATAGTGCT  
TCTCATTTG (SEQ ID NO:1)

FIGURE 1B

underlined = deleted in targeting construct

**bold** = sequence flanking Neo insert in targeting construct

AGCAGAGAGCCTGGTGGGCATGGACATCTTTATCCACATACCTTAGTGTGACCACGCCGA  
 CAGAAAAC**TACTAAGGCCATCTCAGGGGTGCCTGTGCCAGGAGAGGGGGCGGTGTCCCC**  
**GGGCCGCAGAGCCATGCCTTTTCGGCCTGAAGCTCCGCAGGACTCGGCGCTACAACGTCCT**  
**GAGCAAGAACTGCTTTGTTGCCCGGATCCGCCTGCTGGACAGCAATGTCATCGAGTGCAC**  
**GCTGTCGGTGGAAAGCACGGGGCAAGAGTGCCTGGAGGGCCGTGGCCCAGAGGCTGGAGCT**  
**GAGGGAGACGC**ACTACTTCGGCCTTTGGTTTCTCAGCAAGAGCCAGCAGGCGAGATGGGT  
 AGAGCTGGAGAAGCCACTGAAGAAACATCTGGACAAGTTTGCTAACGAGCCTCTGCTTTT  
 CTTCCGGAGTCATGTTCTATGTGCCAAATGTGTACGGCTTCAGCAGGAGGCCACAAGATA  
 TCAGTATTACCTGCAAGTCAAAAAAGACGTGCTTGAAGGACGGTTGCGGTGCTCCCTGGA  
 ACAAGTGATCCGGCTGGCTGGCTTAGCTGTGCAAGCTGACTTCGGAGATTATAACCAAGTT  
 TGATTCCCAAGAGTTCTCCGAGAGTATGTGCTCTTTCCTATGGATTGGCCATGGAGGA  
 GGCGGCTCTGGAGGAGCTAACCCAGAAGGTGGCCAGGAACACAAAGCTCATAGCGGGAT  
 CCTGCCGGCTGAAGCTGAAGTATGATGATCAACGAGGTAGAGCGTTTGGATGGATTGG  
 ACAGGAGATCTTCCCCGTGAAGGACAGTCATGGCAACAGCGTGCACCTCGGCATCTTCTT  
 CATGGGGATTTTTGTGAGGAACAGGGTCGGGAGACAGGCAGTGATATACAGGTGGAATGA  
 CATTTGGGAGTGTTACTCACAGCAAAGCAGCCATCCTGTTGGAGCTGATTGACAAGGAGGA  
 GACCGCGCTCTTCCATACAGATGATATTGAAAATGCCAAGTACATTTCTCGGTTGTTTAC  
 CACTCGGCACAAATTTTACAAACAGAACAAGATCTGCACTGAACAGTCAAATTTCTCCACC  
 CCAATCAGACGCCAGCCACCTGGAGCGGTCCTTACTGCCAAGGCAGCAGCCGTATAT  
 CTTGCCTCCCATGCATGTCCAGTGCAGTGAGCACTACTCGGAGACCCATACTTCCCAAGA  
 CAGCATTTTCCCCGGGAACGAAGAAGCCTTGTAAGTGCCTTCTCACAACAGCCGACCT  
 TAATTACTTGAACGGCACCGTCACCAATGGCAGCGTGTGCAGCGTTCACAGCGTCAACTC  
 CCTCAGCTGCTCCCAGAGCTTCATTACAGGCGTCTCCAGTGTCTCCAACCTTAGCATCCC  
 TGGGAGTGACATCATGAGGGCCGATTACATCCCCAGCCACCGCCACAGCACCATCATCGT  
 GCCGTCTTACAGGCCGACCCCAGATTACGAGACGGTCATGAGGCAGATGAAGAGGGGTCT  
 GATGCACGCAGACAGCCAGAGCCGGTCTCTGCGTAACCTCAATATCATCAACCCCATGC  
 CTATAACCAGCCCCGAGGAAGTGGTGTACAGCCAGCCGGAGATGCGGGAGAGGCATCCCTA  
 CACGGTCCCCCTATGCACACCAGGGGTGCTACGGTCACAACTTGTAAGTCCGTCTGACCA  
 GATGAACCCCCAAAATTGTGCGATGCCTATCAAGCCAGGGGCCAGTTCCATCTCTCACAC  
 AGTGAGCACTCCAGAACTAGCCAACATGCAGCTCCAAGGAGCACAACACTATAGCACAGC  
 CCACATGCTCAAGAACTATCTATTACAGGCCGCCACCCCTTACCCTCGGCCCCGTCTGC  
 CACCAGCACCCAGACCTCGCCAGCCACCGCCACAAGTACGTACGCGGCAGCAGCCCTGA  
 TCTGGTAACTCGGAAGGTGCAGCTCTCCGTAAAGACCTTCCAGGAGGACAGCTCACCTGT  
 GGTCCATCAGTCTCTGCAGGAGGTGAGCGAACCCCTCACAGCCACCAAGCACCATGGCGG  
 CGGCGGTGGCACGGTGAATAAACGCCACAGCCTGGAGGTGATGAACAGCATGGTGAGAGG  
 CATGGAGGCCATGACACTGAAGTCACTCAATATCCCCATGGCTCGCCGCAACACCCTTCG  
 GGAGCAGGGCCCTTCCGAGGACGGGCGGCCACGAAGTGCACGGTCTCCCCAGTATCA  
 CCACAAGAAGACATTCTCGGATGCCACCATGCTGATCCACAGCAGTGAGAGCGAGGAAGA

FIGURE 2A

### FIGURE 2B

Gene Sequence  
Structure \*

192 bp

Sequence Deleted

274 bp

Size of full-length  
cDNA: 3957 bp

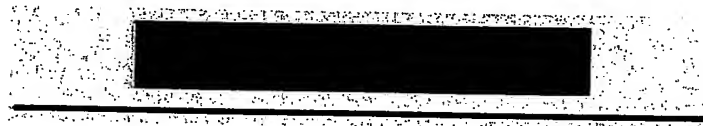
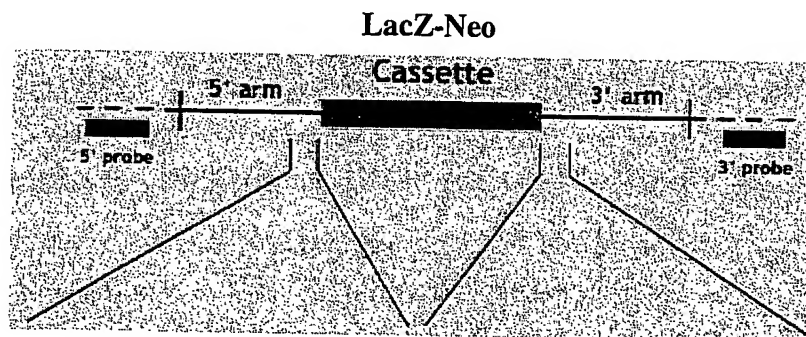


FIGURE 3

Targeting Vector\*  
(genomic sequence)

Arm Length:  
5': 3.5 kb  
3': 2 kb



Targeting Vector  
Endogenous Locus

\* Not drawn to scale

5' >CAGCTGCCCCGGCAGAGAGCCT  
GGTGGGCATGGACATCTTTATCCA  
CATACCTTAGTGTGACCACGCCGA  
CAGAAACTACTAAGGCCATCTCA  
GGGGTGCCCTGTGCCAGGAGAGGGG  
GGCGGTGTCCCCGGGCCGAGAGC  
CATGCCCTTTCGGCCTGAAGCTCCG  
CAGGACTCGGCGCTACAACGTCCT  
GAGCAAGAACT<3' (SEQ ID  
NO: 2)

5' >GAGGCCGTGGCCCAGAGGCTG  
GAGCTGAGGGAGGTGAGTTGAGCG  
CGCATCCCTGCCTGTTGTGTGGAC  
AGGGAGTGGGCTCTTCAGAGGAAC  
CAGCTATCTGCTTAACGTGTTGGC  
ACCTGCTGTGTTTTTCAGCCTAAGC  
GTGTGTTTAAAGAACCTGCTTTT  
CTAGGGTGGGTGTGGCCCCGGGA  
AGTTCCAGCAT<3' (SEQ ID  
NO: 3)

FIGURE 4